

GenCore version 5.1.3
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OM nucleic - nucleic search, using sw model

Run on: October 23, 2002, 06:19:54 ; Search time 983.832 Seconds
(without alignments)
1169.874 Million cell updates/sec

Title: US-09-938-013A-2
Perfect score: 55
Sequence: 1 gaaatgtgcatagacag.....ctaaagaacgacagacag 55

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 1797656 seqs, 10463268293 residues
Total number of hits satisfying chosen parameters: 3595312

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : GenEmbl:*
1: gb_ba:*
2: gb_htg:*
3: gb_in:*
4: gb_om:*
5: gb_ov:*
6: gb_pat:*
7: gb_ph:*
8: gb_pl:*
9: gb_pr:*
10: gb_ro:*
11: gb_sts:*
12: gb_sy:*
13: gb_un:*
14: gb_vi:*
15: em_ba:*
16: em_fun:*
17: em_hum:*
18: em_in:*
19: em_mu:*
20: em_om:*
21: em_or:*
22: em_ov:*
23: em_pat:*
24: em_ph:*
25: em_pl:*
26: em_ro:*
27: em_sts:*
28: em_un:*
29: em_vi:*
30: em_htg_hum:*
31: em_htg_inv:*
32: em_htg_other:*
33: em_htgo_inv:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Match	Length	ID	Description

1	55	100.0	1266	9	HSSMNEUR8	U43883 Human survi
2	55	100.0	1408	6	A77034	A77034 Sequence 11
3	55	100.0	1408	6	A77036	A77036 Sequence 13
4	55	100.0	1408	6	AR100219	AR100219 Sequence
5	55	100.0	1408	6	AR100221	AR100221 Sequence
6	55	100.0	1442	9	BC000908	BC000908 Homo sapi
7	55	100.0	1466	9	HSU21914	U21914 Human dupli
8	55	100.0	1491	9	HSU18423	U18423 Human spina
9	55	100.0	1525	9	BC015308	BC015308 Homo sapi
10	55	100.0	1582	6	A77033	A77033 Sequence 10
11	55	100.0	1582	6	A77035	A77035 Sequence 12
12	55	100.0	1582	6	AR100218	AR100218 Sequence
13	55	100.0	1582	6	AR100220	AR100220 Sequence
14	55	100.0	3271	6	AR100228	AR100228 Sequence
15	55	100.0	82976	9	AC005031	AC005031 Homo sapi
16	55	100.0	113213	9	AC016554	AC016554 Homo sapi
17	55	100.0	129541	9	AC004999	AC004999 Homo sapi
18	55	100.0	131078	9	HSU80017	U80017 Homo sapien
19	55	100.0	155306	2	AC010272	AC010272 Homo sapi
20	55	100.0	155974	9	AC022119	AC022119 Homo sapi
21	55	100.0	168814	9	AC010237	AC010237 Homo sapi
22	55	100.0	190871	9	AC044797	AC044797 Homo sapi
23	46	83.6	885	6	AR022360	AR022360 Sequence
24	32	58.2	1643	9	HSU91640	U91640 Human survi
25	32	58.2	42210	9	AC002050	AC002050 Homo sapi
26	32	58.2	155780	9	HS336012	AL513498 Homo sapi
27	32	58.2	175695	2	AL606844	AL606844 Homo sapi
28	32	58.2	178631	9	AL512635	AL512635 Human DNA
29	30	54.5	206074	9	AC009484	AC009484 Homo sapi
30	28	50.9	113735	2	AC091881	AC091881 Homo sapi
31	28	50.9	138841	2	AC092354	AC092354 Homo sapi
32	28	50.9	156486	2	AC011657	AC011657 Homo sapi
33	28	50.9	163384	9	AC009180	AC009180 Homo sapi
34	27.4	49.8	154157	9	HS101D08	AL133492 Homo sapi
35	27.4	49.8	340000	9	HS21C103	AL163303 Homo sapi
36	27.2	49.5	164988	2	AC080070	AC080070 Homo sapi
37	27.2	49.5	177720	9	AC019193	AC019193 Homo sapi
38	27	49.1	149546	9	AC087256	AC087256 Homo sapi
39	27	49.1	152167	2	AC073307	AC073307 Homo sapi
40	27	49.1	168227	9	AC021868	AC021868 Homo sapi
41	27	49.1	173239	9	AC006478	AC006478 Homo sapi
42	27	49.1	196817	9	AC019209	AC019209 Homo sapi
43	26.8	48.7	259487	10	AL365322	AL365322 Mouse DNA
44	26.2	47.6	111949	2	AL353755	AL353755 Homo sapi
45	26.2	47.6	212854	2	AC093028	AC093028 Homo sapi

ALIGNMENTS

RESULT 1	HSSMNEUR8	1266 bp	DNA	linear	PRI 16-MAY-1996
LOCUS	Human survival motor neuron (SMN) gene, exons 7 and 8, and complete cds.				
DEFINITION	U43883				
ACCESSION	U43883.1	GI:1314344			
VERSION	spinal muscular atrophy gene.				
KEYWORDS	8 of 8				
SEGMENT	human.				
SOURCE	Homo sapiens				
ORGANISM	Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.				
REFERENCE	1 (bases 1 to 1266)				
AUTHORS	Lefebvre,S., Burglen,L., Reboullet,S., Clermont,O., Bulet,P., Viollet,L., Benichou,B., Cruaud,C., Millasseau,P., Zeviani,M. et al.				
TITLE	Identification and characterization of a spinal muscular atrophy-determining gene				
JOURNAL	Cell 80 (1), 155-165 (1995)				
MEDLINE	95112343				
REFERENCE	2 (bases 1 to 1266)				
AUTHORS	Burglen,L., Lefebvre,S., Clermont,O., Bulet,P., Viollet,L.,				

*TITLE Cruaud,C., Munnich,A. and Melki,J.
Structure and organization of the human survival motor neurone
(SMN) gene
JOURNAL Genomics 32 (3), 479-482 (1996)
MEDLINE 96435930
REFERENCE 3 (bases 1 to 1266)
AUTHORS Burglen,I.
TITLE Direct Submission
JOURNAL Submitted (26-DEC-1995) Lydie Burglen, INSERM U-393, IFRM, Hopital
Necker-Enfants Malades, 149, rue de Sevres Cedex 15, Paris 75743,
France
COMMENT Duplicated gene within the 5q13 region; interspecies conserved
sequences. The SMN gene is present in two copies within the SMA
(spinal muscular atrophy) candidate region on chromosome 5q13. The
two copies are transcribed and are distinguished by sequence
differences specific to the centromeric copy.
FEATURES
Location/Qualifiers
1..1266
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/chromosome="5"
/map="5q13"
join(U43876.1:575..688,U43877.1:104..175,
U43878.1:118..237,U43879.1:84..284,U43880.1:69..221,
U43881.1:103..198,U43882.1:53..163,209..262,707..>1266)
/gene="SMN"
/note="spinal muscular atrophy gene; cDNA sequence found
in GenBank Accession Number U18423"
/product="survival motor neuron"
join(U43876.1:575..762,U43877.1:1..257,U43878.1:1..314,
U43879.1:1..357,U43880.1:1..302,U43881.1:1..251,
U43882.1:1..244,1..1266)
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U43881.1:103..198,U43882.1:53..163,209..259)
/gene="SMN"
/note="spinal muscular atrophy gene"
/codon_start=1
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/protein_id="AAC50473.1"
/db_xref="GI:1314346"
/translation="MAMSSGGSGGGVPEQEDSVLFRRTGSDSDIWDITLIKAYD
KAVAFKALKNGDICTSGKPKTPKPKPAKKNKSKKNTAASLQWKVGDKCSAIW
SEDCIYPATIASIDFKETCVVYTYGNREONLSDLLSPICEVANNIEQNAQENE
NESQVSTDESENRSFGKNSDNIRKPSAPWNSFLPPPPMPGRLGPKGLKFNQPP
PPPPPPHLLSCWLPFPFSGPPIIPPPPPICPDSLDADALGSLISWYMSGYHTGY
YMGFRQKQEGRCSHSLN"
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/number=6
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/note="a nucleotide change specific to the centromeric
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/citation=[1]
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209..262
/gene="SMN"
/number=7
misc_difference 214
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/note="a nucleotide change specific to the centromeric
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/replace="t"
join(260..>262,707..>1266)
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263..706
/gene="SMN"
/number=7
misc_difference 362
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/note="a nucleotide change specific to the centromeric
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/citation=[1]
/replace="g"
misc_difference 477
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/note="a nucleotide change specific to the centromeric
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/citation=[1]
/replace="g"
707..>1266
/gene="SMN"
/number=8
polyA_signal 1253..1258
/gene="SMN"
BASE COUNT 430 a 166 c 245 g 425 t
ORIGIN
Query Match 100.0%; Score 55; DB 9; Length 1266;
Best Local Similarity 100.0%; Pred. No. 7.1e-09;
Matches 55; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 GAAATGCTGGCATAGACGACGACCTAAATGACACCACTAAAGAAACGATCAGACAG 55
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Db 707 GAAATGCTGGCATAGACGACGACCTAAATGACACCACTAAAGAAACGATCAGACAG 761
|||||
RESULT 2
A77034
LOCUS A77034 1408 bp DNA linear PAT 19-OCT-1999
DEFINITION Sequence 11 from Patent EP0708178.
ACCESSION A77034
VERSION A77034.1 GI:6088824
KEYWORDS
SOURCE unidentified.
ORGANISM unidentified
unclassified.
REFERENCE 1 (bases 1 to 1408)
AUTHORS Melki,J. and Munnich,A.
TITLE SURVIVAL MOTOR NEURON (SMN) GENE: A GENE FOR SPINAL MUSCULAR
ATROPHY
JOURNAL Patent: EP 0708178-A 11 24-APR-1996;
INST NAT SANTE RECH MED (FR)
FEATURES
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1..1408
/organism="unidentified"
/db_xref="taxon:32644"
BASE COUNT 464 a 198 c 277 g 469 t
ORIGIN
Query Match 100.0%; Score 55; DB 6; Length 1408;
Best Local Similarity 100.0%; Pred. No. 7e-09;
Matches 55; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 GAAATGCTGGCATAGACGACGACCTAAATGACACCACTAAAGAAACGATCAGACAG 55
|||||
Db 846 GAAATGCTGGCATAGACGACGACCTAAATGACACCACTAAAGAAACGATCAGACAG 900
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RESULT 3
A77036
LOCUS A77036 1408 bp DNA linear PAT 19-OCT-1999
DEFINITION Sequence 13 from Patent EP0708178.
ACCESSION A77036
VERSION A77036.1 GI:6088826
KEYWORDS
SOURCE unidentified.
ORGANISM unidentified
unclassified.
REFERENCE 1 (bases 1 to 1408)
AUTHORS Melki,J. and Munnich,A.
TITLE SURVIVAL MOTOR NEURON (SMN) GENE: A GENE FOR SPINAL MUSCULAR
ATROPHY